Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: Figure 1 f: Monte Carlo simulation results for the 3-bed EVMAP.

File Name: Supplementary Data 2

Description: Figure 1 g: Calculated sample processing time for 3-bed (5μL/min) and 7-bed

devices (20 µL/min).

File Name: Supplementary Data 3

Description: Figure 1 h: Results of Monte Carlo simulation for the 7-bed EVMAP

File Name: Supplementary Data 4

Description: Figure 2 h: NTA results and TEM images showing the number of EVs released

during first and second USER enzyme release.

File Name: Supplementary Data 5

Description: Figure 2k: Percentage of EVs released during first and second release with

USER enzyme.

File Name: Supplementary Data 6

Description: Figure 3a: Cell line viability when cultured with different LPS concentrations in

culture medium.

File Name: Supplementary Data 7

Description: Figure 3c: mRNA gene expression profiles for CD8+ EVs cells (0.5 and 0.7 ng of RNA for stimulated and unstimulated respectively used in RT reactions). Figure 3d: mRNA gene expression profiles for CD8+ Molt-3 cells (0.7 ng RNA for stimulated and unstimulated was used in RT reactions). Figure 3e: Correlation between mRNA copies found in stimulated and unstimulated Molt-3 cells and EVs. Figure 3f: Correlation between mRNA copies found in EVs and Molt-3 cells in stimulated and unstimulated conditions.

File Name: Supplementary Data 8

Description: Figure 4d: Correlation plot of particle concentration (presumably EVs) with RNA mass isolated from affinity selected CD8+ EVs.

File Name: Supplementary Data 9

Description: Figure 4e: Boxplots comparing the gene expression of CD8+Tcells and CD8+EVs isolated from healthy donor plasma for PLBD1 for cells (n=5) and EVs (n=6). Figure 4f: Boxplots comparing the gene expression of CD8+T-cells and CD8+EVs isolated from healthy donor plasma for vFOS for cells (n=5) and EVs (n=6). Figure 4g: Boxplots comparing the gene expression of CD8+T-cells and CD8+EVs isolated from healthy donor plasma for MMP9 for cells (n=5) and EVs (n=6). Figure 4h: Boxplots comparing the gene expression of CD8+T-cells and CD8+EVs isolated from healthy donor plasma for CA4 for cells (n=5) and EVs (n=6). Figure 4i: Boxplots comparing the gene expression of CD8+T-cells and CD8+EVs isolated from healthy donor plasma for VCAN for cells (n=5) and EVs (n=6).

File Name: Supplementary Data 10

Description: Figure 5a: NTA images of EVs isolated from clinical sample #4 by PEG precipitation and affinity selected with anti-CD8 α mAb uisng the 7-bed EV-MAP.

File Name: Supplementary Data 11

Description: Figure 5e: Heat maps presenting the EV mRNA expression profiles for sample

#4.

File Name: Supplementary Data 12

Description: Figure 5f: NTA results for selected Samples 1,4,6,and 8.

File Name: Supplementary Data 13

Description: Figure 5g: mRNA expression profiling for selected genes in clinical samples. Figure 5h: Heat map analysis of clinical samples (marked with numbers) and healthy donors

(identified with letters). Figure 5i: Principal component analysis for clinical samples

(identified with numbers) and healthy donors (identified with letters).